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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/045,992

DATE: 01/28/2002
TIME: 14:14:39

Input Set : A:\053689-5006-01.ST25.txt
Output Set: N:\CRF3\01282002\J045992.raw

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2 <110> APPLICANT: LINDNER, Volkhard
3 FRIESEL, Robert E.
5 <120> TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO REMODELIN
7 <130> FILE REFERENCE: 053689-5006-01
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/045,992
C--> 9 <141> CURRENT FILING DATE: 2001-10-19
9 <150> PRIOR APPLICATION NUMBER: US 09/692,081
10 <151> PRIOR FILING DATE: 2000-10-19
12 <160> NUMBER OF SEQ ID NOS: 9
14 <170> SOFTWARE: PatentIn version 3.1
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 1192
18 <212> TYPE: DNA
19 <213> ORGANISM: Rattus sp.
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23 tgcctcctgc tctgcgcttc gcagctaccg cacacgatgc acccccaagg ccgcgccgcc 120
24 tccccacagc tgctgctcgg cctcttcctt gtgctactgc tgcttctgca gctgtccgcg 180
25 ccgtccagcg cctctgagaa tcccaaggtg aagcaaaaag cgctgatccg gcagagggaa 240
26 gtggtagacc tgtataatgg gatgtgccta caaggaccag caggagttcc tggtcgcgat 300
27 gggagccctg gggccaatgg cattcctggc acaccgggaa tcccaggtcg ggatggattc 360
28 aaaggagaga aaggggagtg cttaagggaa agctttgagg aatcctggac cccaaactac 420
29 aagcagtgtt catggagttc acttaattat ggcatagatc ttgggaaaat tgcggaatgt 480
30 acattcacaa agatgcgatc caacagcgct cttcgagttc tgttcagtgg ctcgcttcgg 540
31 ctcaaatgca ggaatgcttg ctgtcaacgc tggtatttta cctttaatgg agctgaatgt 600
32 tcaggacctc ttcccattga agctatcatc tatctggacc aaggaagccc tgagttaaat 660
33 tcaactatta atattcatcg tacttcctcc gtggaaggac tctgtgaagg gattggtgct 720
34 ggactggtag acgtggccat ctgggtcggc acctgttcag attaccccaa aggagacgct 780
35 tctactgggt ggaattctgt gtcccgcatc atcattgaag aactaccaaa ataaagcccc 840
36 tgaaggtttc attccctgcc tcatttactt gttaaatcaa gcctctggat gggtcattta 900
37 aatgacattt cagaagtcac ttatgtgctc agccaaatga aaaagcaaag ttaaatacgt 960
38 ttacagacca aagtgtgatc tcacacttta agatctagca ttatccattt tatttcaacc 1020
39 aaagatggtt tcaggatttt atttctcatt gattactttt tgagcctata taccggaatg 1080
40 ctgttatagt ctttaatatt tcctactgtt gacattttga aacatataaa agttatgtct 1140
41 ttgtaagagc tgtatagaat tattttatat gttaaataaa tgcttcaaac aa 1192
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44 <211> LENGTH: 245
45 <212> TYPE: PRT
46 <213> ORGANISM: Rattus sp.
48 <400> SEQUENCE: 2
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51 1 5 10 15
53 Phe Leu Val Leu Leu Leu Leu Leu Gln Leu Ser Ala Pro Ser Ser Ala

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56 Ser Glu Asn Pro Lys Val Lys Gln Lys Ala Leu Ile Arg Gln Arg Glu
57          35                  40                  45
59 Val Val Asp Leu Tyr Asn Gly Met Cys Leu Gln Gly Pro Ala Gly Val
60      50                  55                  60
62 Pro Gly Arg Asp Gly Ser Pro Gly Ala Asn Gly Ile Pro Gly Thr Pro
63 65                  70                  75                  80
65 Gly Ile Pro Gly Arg Asp Gly Phe Lys Gly Glu Lys Gly Glu Cys Leu
66                  85                  90                  95
68 Arg Glu Ser Phe Glu Glu Ser Trp Thr Pro Asn Tyr Lys Gln Cys Ser
69              100                 105                 110
71 Trp Ser Ser Leu Asn Tyr Gly Ile Asp Leu Gly Lys Ile Ala Glu Cys
72          115                 120                 125
74 Thr Phe Thr Lys Met Arg Ser Asn Ser Ala Leu Arg Val Leu Phe Ser
75      130                 135                 140
77 Gly Ser Leu Arg Leu Lys Cys Arg Asn Ala Cys Cys Gln Arg Trp Tyr
78 145                 150                 155                 160
80 Phe Thr Phe Asn Gly Ala Glu Cys Ser Gly Pro Leu Pro Ile Glu Ala
81                  165                 170                 175
83 Ile Ile Tyr Leu Asp Gln Gly Ser Pro Glu Leu Asn Ser Thr Ile Asn
84              180                 185                 190
86 Ile His Arg Thr Ser Ser Val Glu Gly Leu Cys Glu Gly Ile Gly Ala
87          195                 200                 205
89 Gly Leu Val Asp Val Ala Ile Trp Val Gly Thr Cys Ser Asp Tyr Pro
90      210                 215                 220
92 Lys Gly Asp Ala Ser Thr Gly Trp Asn Ser Val Ser Arg Ile Ile Ile
93 225                 230                 235                 240
95 Glu Glu Leu Pro Lys
96                  245
98 <210> SEQ ID NO: 3
99 <211> LENGTH: 1220
100 <212> TYPE: DNA
101 <213> ORGANISM: Homo sapiens
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105 cctccgcctc cagctccgcg ctgcccggca gccgggagcc atgcgacccc agggccccgc    120
106 cgcctccccg cagcggctcc gcggcctcct gctgctcctg ctgctgcagc tgcccgcgcc    180
107 gtcgagcgcc tctgagatcc ccaaggggaa gcaaaaggcg cagctccggc agagggaggt    240
108 ggtggacctg tataatggaa tgtgcttaca agggccagca ggagtgcctg gtcgagacgg    300
109 gagccctggg gccaatggca ttccgggtac acctgggatc ccaggtcggg atggattcaa    360
110 aggagaaaag ggggaatgtc tgagggaaag ctttgaggag tcctggacac ccaactacaa    420
111 gcagtgttca tggagttcat tgaattatgg catagatctt gggaaaattg cggagtgtac    480
112 atttacaaag atgcgttcaa atagtgctct aagagttttg ttcagtggct cacttcggct    540
113 aaaatgcaga aatgcatgct gtcagcgttg gtatttcaca ttcaatggag ctgaatgttc    600
114 aggacctctt cccattgaag ctataattta tttggaccaa ggaagccctg aaatgaattc    660
115 aacaattaat attcatcgca cttcttctgt ggaaggactt tgtgaaggaa ttggtgctgg    720
116 attagtggat gttgctatct gggttggcac ttgttcagat tacccaaaag gagatgcttc    780
117 tactggatgg aattcagttt ctcgcatcat tattgaagaa ctaccaaaat aaatgcttta    840
118 attttcattt gctacctctt tttttattat gccttggaat ggttcactta aatgacattt    900
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119 taaataagtt tatgtataca tctgaatgaa aagcaaagct aaatatgttt acagaccaaa    960
120 gtgtgatttc acactgtttt taaatctagc attattcatt ttgcttcaat caaaagtggt   1020
121 ttcaatattt ttttagttgg ttagaatact ttcttcatag tcacattctc tcaacctata   1080
122 atttggaata ttgttgtggt cttttgtttt ttctcttagt atagcatttt taaaaaaata   1140
123 taaaagctac caatctttgt acaatttgta aatgttaaga attttttttta tatctgttaa   1200
124 ataaaaatta tttccaacaa                                               1220
126 <210> SEQ ID NO: 4
127 <211> LENGTH: 243
128 <212> TYPE: PRT
129 <213> ORGANISM: Homo sapiens
131 <400> SEQUENCE: 4
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136 Leu Leu Leu Leu Leu Leu Gln Leu Pro Ala Pro Ser Ser Ala Ser Glu
137             20                  25                  30
139 Ile Pro Lys Gly Lys Gln Lys Ala Gln Leu Arg Gln Arg Glu Val Val
140         35                  40                  45
142 Asp Leu Tyr Asn Gly Met Cys Leu Gln Gly Pro Ala Gly Val Pro Gly
143     50                  55                  60
145 Arg Asp Gly Ser Pro Gly Ala Asn Gly Ile Pro Gly Thr Pro Gly Ile
146 65                  70                  75                  80
148 Pro Gly Arg Asp Gly Phe Lys Gly Glu Lys Gly Glu Cys Leu Arg Glu
149                 85                  90                  95
151 Ser Phe Glu Glu Ser Trp Thr Pro Asn Tyr Lys Gln Cys Ser Trp Ser
152             100                 105                 110
154 Ser Leu Asn Tyr Gly Ile Asp Leu Gly Lys Ile Ala Glu Cys Thr Phe
155         115                 120                 125
157 Thr Lys Met Arg Ser Asn Ser Ala Leu Arg Val Leu Phe Ser Gly Ser
158     130                 135                 140
160 Leu Arg Leu Lys Cys Arg Asn Ala Cys Cys Gln Arg Trp Tyr Phe Thr
161 145                 150                 155                 160
163 Phe Asn Gly Ala Glu Cys Ser Gly Pro Leu Pro Ile Glu Ala Ile Ile
164                 165                 170                 175
166 Tyr Leu Asp Gln Gly Ser Pro Glu Met Asn Ser Thr Ile Asn Ile His
167             180                 185                 190
169 Arg Thr Ser Ser Val Glu Gly Leu Cys Glu Gly Ile Gly Ala Gly Leu
170         195                 200                 205
172 Val Asp Val Ala Ile Trp Val Gly Thr Cys Ser Asp Tyr Pro Lys Gly
173     210                 215                 220
175 Asp Ala Ser Thr Gly Trp Asn Ser Val Ser Arg Ile Ile Ile Glu Glu
176 225                 230                 235                 240
178 Leu Pro Lys
181 <210> SEQ ID NO: 5
182 <211> LENGTH: 277
183 <212> TYPE: PRT
184 <213> ORGANISM: Rattus sp.
186 <400> SEQUENCE: 5
188 Met Arg Pro Ala Ala Glu Leu Gly Gln Thr Leu Ser Arg Ala Gly Leu
189 1               5                  10                  15
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191 Cys Arg Pro Leu Cys Leu Leu Leu Cys Ala Ser Gln Leu Pro His Thr
192             20                  25                  30
194 Met His Pro Gln Gly Arg Ala Ala Ser Pro Gln Leu Leu Leu Gly Leu
195         35                  40                  45
197 Phe Leu Val Leu Leu Leu Leu Leu Gln Leu Ser Ala Pro Ser Ser Ala
198     50                  55                  60
200 Ser Glu Asn Pro Lys Val Lys Gln Lys Ala Leu Ile Arg Gln Arg Glu
201 65                  70                  75                  80
203 Val Val Asp Leu Tyr Asn Gly Met Cys Leu Gln Gly Pro Ala Gly Val
204                 85                  90                  95
206 Pro Gly Arg Asp Gly Ser Pro Gly Ala Asn Gly Ile Pro Gly Thr Pro
207             100                 105                 110
209 Gly Ile Pro Gly Arg Asp Gly Phe Lys Gly Glu Lys Gly Glu Cys Leu
210         115                 120                 125
212 Arg Glu Ser Phe Glu Glu Ser Trp Thr Pro Asn Tyr Lys Gln Cys Ser
213     130                 135                 140
215 Trp Ser Ser Leu Asn Tyr Gly Ile Asp Leu Gly Lys Ile Ala Glu Cys
216 145                 150                 155                 160
218 Thr Phe Thr Lys Met Arg Ser Asn Ser Ala Leu Arg Val Leu Phe Ser
219                 165                 170                 175
221 Gly Ser Leu Arg Leu Lys Cys Arg Asn Ala Cys Cys Gln Arg Trp Tyr
222             180                 185                 190
224 Phe Thr Phe Asn Gly Ala Glu Cys Ser Gly Pro Leu Pro Ile Glu Ala
225         195                 200                 205
227 Ile Ile Tyr Leu Asp Gln Gly Ser Pro Glu Leu Asn Ser Thr Ile Asn
228     210                 215                 220
230 Ile His Arg Thr Ser Ser Val Glu Gly Leu Cys Glu Gly Ile Gly Ala
231 225                 230                 235                 240
233 Gly Leu Val Asp Val Ala Ile Trp Val Gly Thr Cys Ser Asp Tyr Pro
234                 245                 250                 255
236 Lys Gly Asp Ala Ser Thr Gly Trp Asn Ser Val Ser Arg Ile Ile Ile
237             260                 265                 270
239 Glu Glu Leu Pro Lys
240         275
242 <210> SEQ ID NO: 6
243 <211> LENGTH: 403
244 <212> TYPE: RNA
245 <213> ORGANISM: Artificial Sequence
247 <220> FEATURE:
248 <223> OTHER INFORMATION: Description of Artificial Sequence:REMODELIN antisense
ribonucleoprobe
250 <400> SEQUENCE: 6
251 ccacccagua gaagcgucuc cuuuggggua aucugaacag gugccgaccc agauggccac     60
252 gucuaccagu ccagcaccaa ucccuucaca gaguccuucc acggaggaag uacgaugaau    120
253 auuaauaguu gaauuuaacu cagggcuucc uugguccaga uagaugauag cuucaauggg    180
254 aagaggуccu gaacauucag cuccauuaaa gguaaaauac cagcguugac agcaagcauu    240
255 ccugcauuug agccgaagcg agccacugaa cagaacucga agagcgcugu uggaucgcau    300
256 cuuugugaau guacauuccg caauuuuccc aagaucuaug ccauaauuaa gugaacucca    360
257 ugaacacugc uuguaguuug ggguccagga uuccucaaag cuu                      403
259 <210> SEQ ID NO: 7
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260 <211> LENGTH: 15
261 <212> TYPE: PRT
262 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: Description of Artificial Sequence:carboxy-terminal amino acids of REMODELIN
267 <400> SEQUENCE: 7
269 Gly Trp Asn Ser Val Ser Arg Ile Ile Ile Glu Glu Leu Pro Lys
270 1 5 10 15
272 <210> SEQ ID NO: 8
273 <211> LENGTH: 24
274 <212> TYPE: PRT
275 <213> ORGANISM: Artificial Sequence
277 <220> FEATURE:
278 <223> OTHER INFORMATION: Description of Artificial Sequence:insulin signal peptide
280 <400> SEQUENCE: 8
282 Met Ala Leu Leu Val His Phe Leu Pro Leu Leu Ala Leu Leu Ala Leu
283 1 5 10 15
285 Trp Glu Pro Lys Pro Thr Gln Ala
286 20
288 <210> SEQ ID NO: 9
289 <211> LENGTH: 734
290 <212> TYPE: DNA
291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <223> OTHER INFORMATION: Description of Artificial Sequence:myc-tagged REMODELIN construct
296 <400> SEQUENCE: 9
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298 tgctgcttct gcagctgtcc gcgccgtcca gcgcctctga gaatcccaag gtgaagcaaa 120
299 aagcgctgat ccggcagagg gaagtggtag acctgtataa tgggatgtgc ctacaaggac 180
300 cagcaggagt tcctggtcgc gatgggagcc ctggggccaa tggcattcct ggcacaccgg 240
301 gaatcccagg tcgggatgga ttcaaaggag agaaagggga gtgcttaagg gaaagctttg 300
302 aggaatcctg gaccccaaac tacaagcagt gttcatggag ttcacttaat tatggcatag 360
303 atcttgggaa aattgcggaa tgtacattca caaagatgcg atccaacagc gctcttcgag 420
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307 gactctgtga agggattggt gctggactgg tagacgtggc catctgggtc ggcacctgtt 660
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VERIFICATION SUMMARY DATE: 01/28/2002
PATENT APPLICATION: US/10/045,992 TIME: 14:14:40

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date